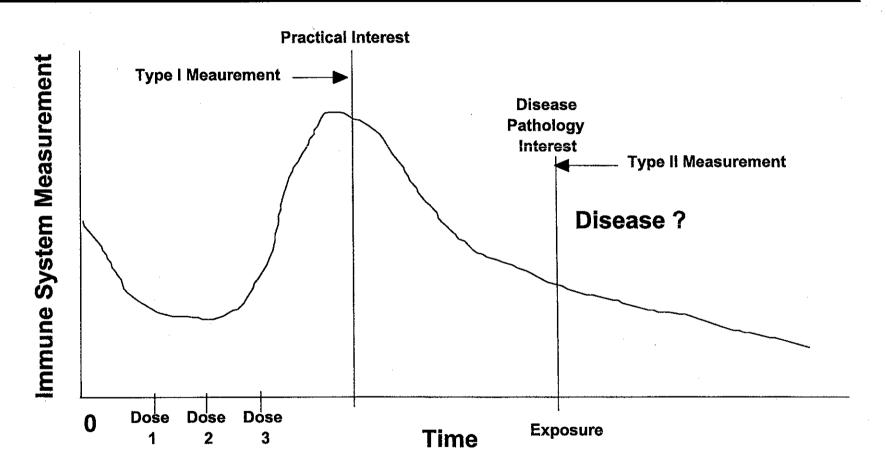
### Vaccine Correlates of Immunity

Robert C. Kohberger, Ph.D. Wyeth-Lederle Vaccines and Pediatrics

### Typical Vaccine Trial



#### Immune System Measurements

#### ◆ Humoral immunity

- based on antibodies which are products of the B cell system
- antibodies found in body fluids (plasma, lymph, and others)
- circulating antibody: concentrations measured in plasma

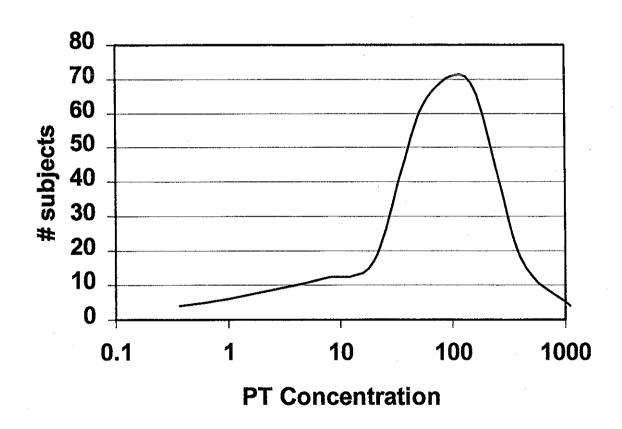
#### ◆ Cell mediated immunity

product of the T cell system

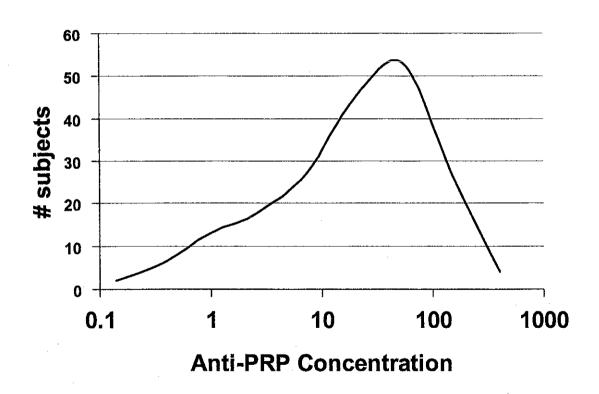
### Vaccine Immunogenicity Trials

- Antibody concentrations measured shortly after completion of the primary dose series (Type I measurement)
- Antibody concentrations measured immediately prior to infection (Type II measurement)
- Antibody concentrations measured at various times after the completion of the primary series (kinetic studies)

## Type I Example: Pertussis Toxin (PT) GMT = 101.8, std dev = 1.6



#### Type I Example: Hib (anti-PRP) GMT = 30.9, std dev = 1.67



### Vaccine Efficacy Trials

- Vaccines are intended to prevent the clinical manifestation of diseases
- Efficacy is determined by identifying cases of disease after vaccination and comparing to cases in a placebo group
- Vaccine efficacy is expressed as a reduction in the relative risk of disease

## Correlates of Immunity (Protection) Uses

- ◆ 'Surrogate' for efficacy
  - allow use of immunogenicity trials (Type I measurements) rather than efficacy trials
- Allow for comparison between different manufacturing processes including combination vaccines
- Allow for prediction of how different antibody response distributions change efficacy

# Correlates of Immunity Common Approaches

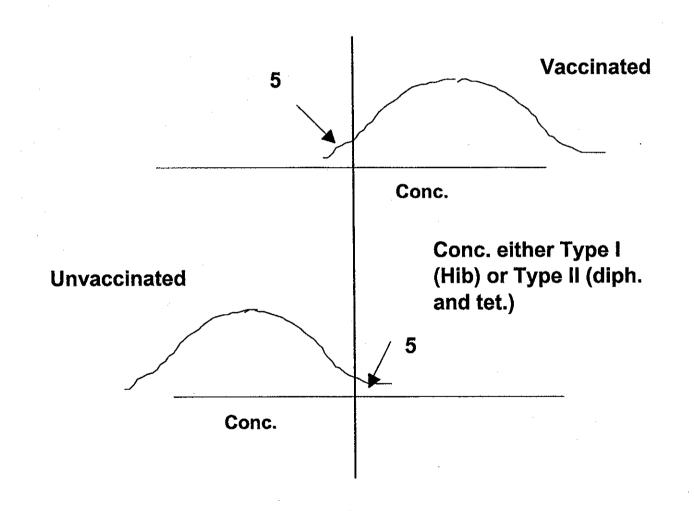
◆ Population based - H. influenza, diphtheria, tetanus

◆ Concentration specific

# Historical Model Population Based

- Measure antibody levels in vaccinated and unvaccinated groups
- Protective level is that which is exceeded by most of the vaccinated group and not reached by most of the unvaccinated group
- ◆ Problem: arbitrary

### Historical Model Population Based



## Concentration Specific Model Logistic Predictive Model of Disease

Probability of disease given a concentration and exposed =  $p(D|c,E) = y = e^{a+bc}/1+e^{a+bc}$ 

where a,b are constants to be estimated c = concentration

y = probability between 0.0 and 1.0

a = log odds of disease when t = 0

b = change in log odds of disease with unit change in concentration

### Vaccine Efficacy

VE = 1 - <u>disease rate given treated</u> disease rate given not treated

P(D|T) = prob (disease given treated) P(D|NT) = prob (disease given not treated)

$$VE = 1 - \frac{P(D|T)}{P(D|NT)}$$

#### Vaccine Efficacy

```
VE = 1 - \underline{P(D|T,E)P(E|T)}
P(D|NT,E)P(E|NT)
since in a randomized, double-blind trial P(E|T) = P(E|NT) \text{ then }
VE = 1 - \underline{P(D|T,E)}
P(D|NT,E)
now, let c = \text{concentration}
p(D|c,E) = e^{a+bc}/(1 + e^{a+bc}) = \text{logistic model}
```

### Vaccine Efficacy

$$VE = 1 - \frac{\int p(D|c,E)f(c|T)dc}{\int p(D|c,E)f(c|NT)dc}$$

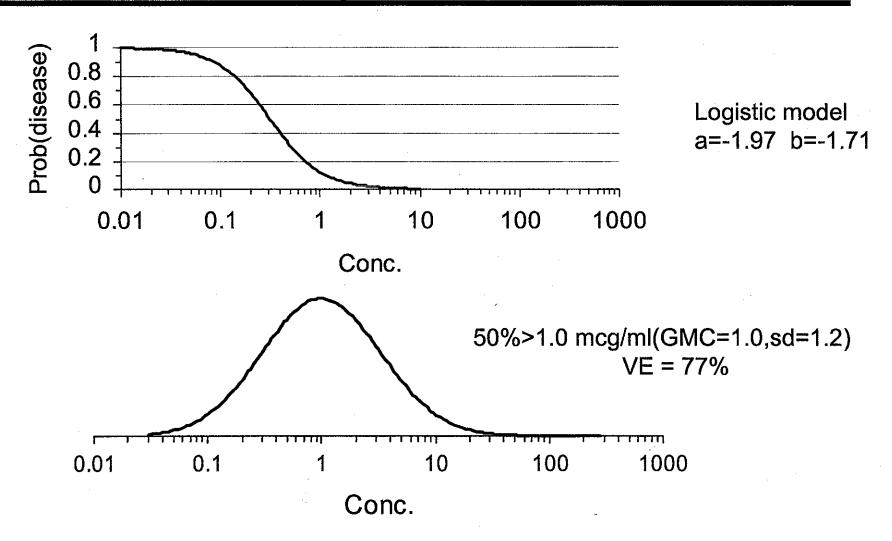
$$=1-\frac{\mathsf{E}_{\mathsf{T}}[\mathsf{p}(\mathsf{D}|\mathsf{c},\mathsf{E})]}{\mathsf{E}_{\mathsf{NT}}[\mathsf{p}(\mathsf{D}|\mathsf{c},\mathsf{E})]}$$

### Vaccine Efficacy Estimate

$$VE = \frac{\sum_{j} \frac{e^{a+bc_{j}}}{1+e^{a+bc_{j}}}}{\frac{e^{a}}{1+e^{a}}}$$

j = 1, ..., n n = number of subjects in immunogenicity trial $c_j = conc.$  in the j-th subject of the trial

## Correlates-Immunogenicity Vaccine Efficacy



# Theoretical Issues Concentration Specific Model

 Accounting for the time-varying nature of the disease risk due, possibly, to the timevarying nature of the immunologic measurement

◆ Accounting for the follow-up time

### Extending the Concentration Specific Model Logit Model

```
In (logit) = a + b^* ln(conc)

= a + b^*(c_0 + f^*time)

c_0 = ln(initial concentration)

(assuming exponential decay of immunologic measurement)
```

- for those without disease time is censored
- Cox regression applications ?

#### Practical Issues

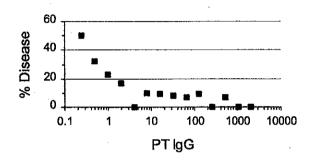
- Limited number of disease cases
- Limited amount of immunology information
- ◆ Level of noise in logit estimation is high considering often only 10-15% of population is exposed to disease
- Differing antibody decay characteristics between subjects

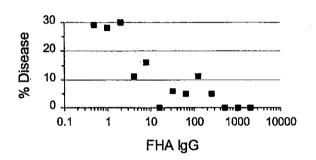
### Example: Pertussis Trials

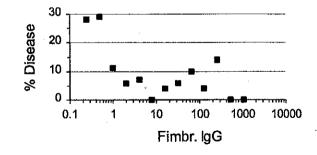
- Cherry, et. al, <u>Vaccine</u>, 1998, 16:1901-1906
   Storsaeter, et. al., <u>Vaccine</u>, 1998, 16:1907-1916
- Use pre-exposure measurement (Type II) imputed - and simple logistic model
- ◆ Do not back estimate to Type I post vaccination measurement

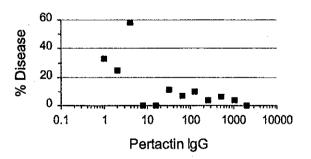
#### Erlangen Acellular Pertussis Trial

Household Contact Sample (subjects exposed in their household) % Disease vs. Antibody Conc. after primary series



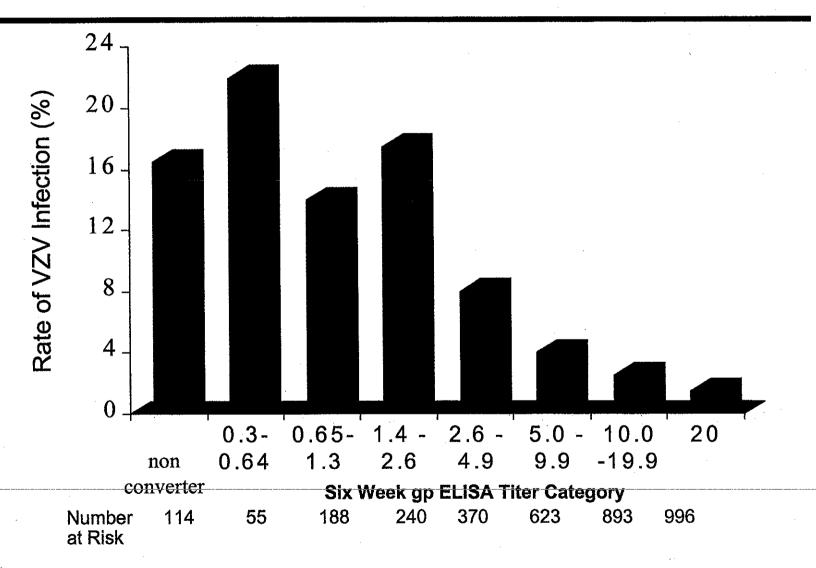






#### Rate of Varicella\* After VZV Vaccine

Adapted from: CJ White et al PIDJ 11:19-23, 1992



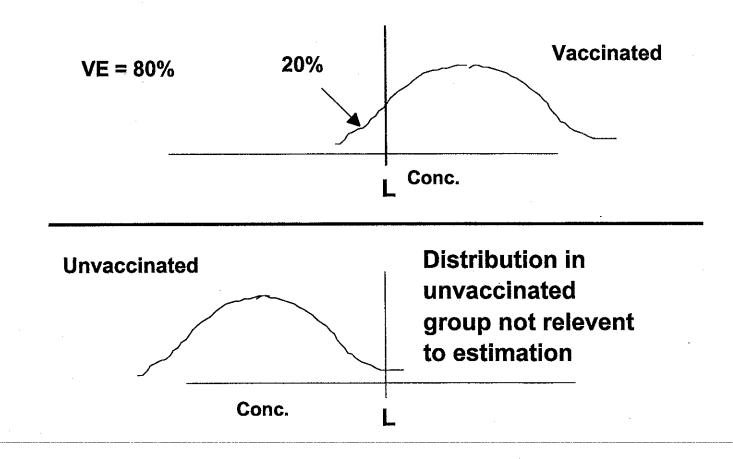
## Population Based Model Is there a Justification?

```
Prob (disease) = 0 if measurement ≥ L
b if measurement < L

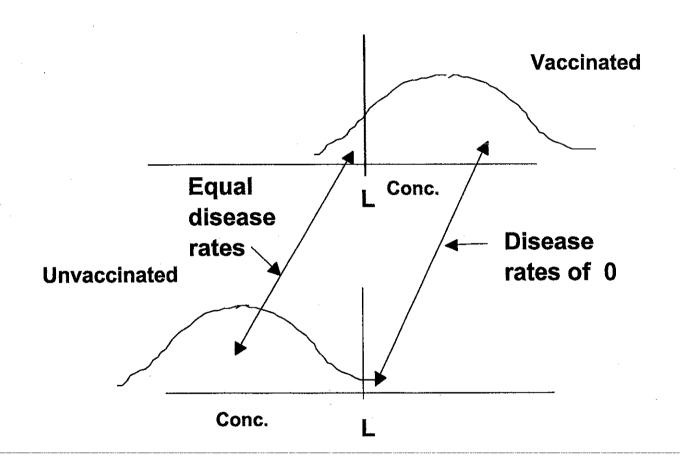
↓
```

VE = % of population with measurement ≥ L

#### Population Based Model



## Population Based Model Model Check



### Basic Immunology Issues Serology IgG Measurements

# , J ( ) 3 ( ).

- ◆ Relationship of assay measurement with opsonic or virus neutralizing capability (functional antibody)
- Amount and speed of anamnestic response
- ◆ Mucosal and cellular immunity

#### Conclusions

◆ Concentration specific models may be too impractical for common use

 Population based models may provide a sufficient approximation